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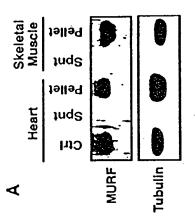


FIG. 3

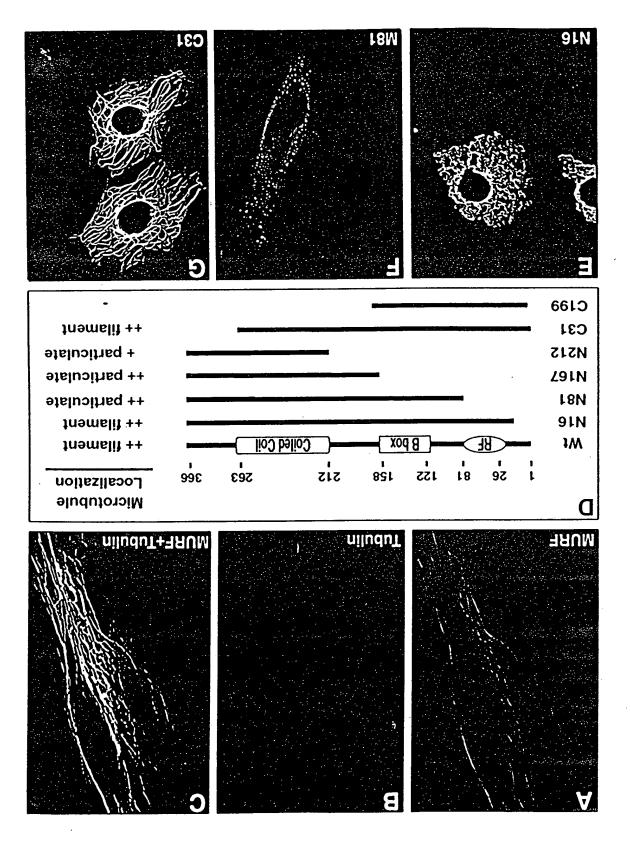
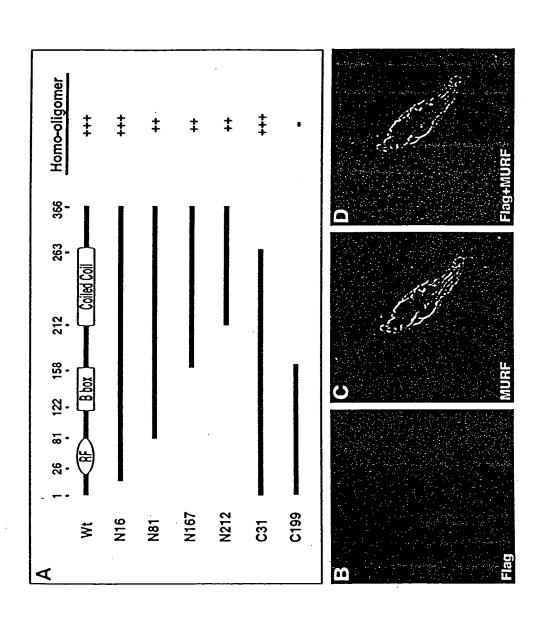


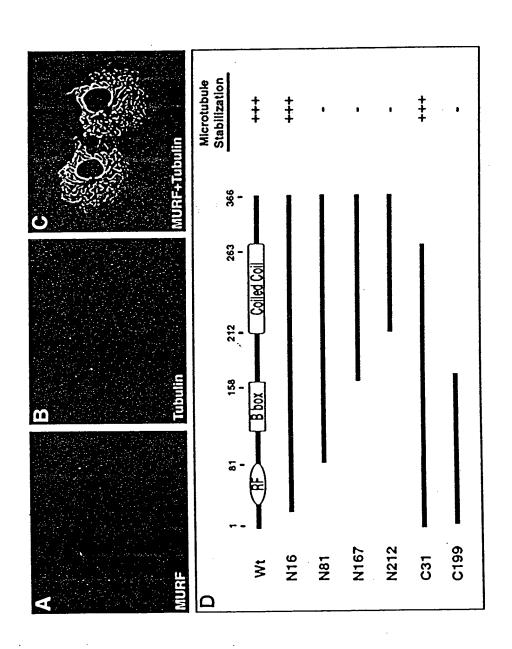
FIG. 4

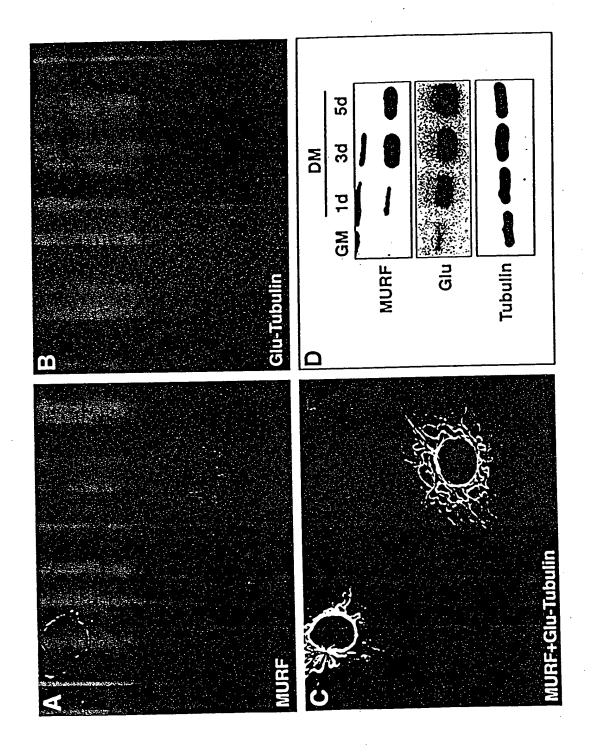
FIG. 5











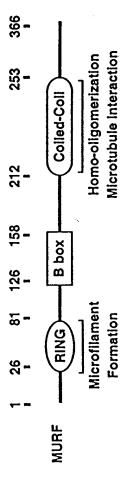


FIG. 9

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1 MNFTVGFKPLLG-DAHNMDNLEKQLICPICLEMFSKFVVILPCQHNLCRK 49
1 MSTSLNYKSFSK-EQQTMDNLEKQLICPICLEMFTKPVVILPCQHNLCRK 49
1 MDYKSSLIPDGNAMENLEKQLICPICLEMFTKPVVILPCQHNLCRK 46
 MURF 1
 MURF 2
 MURF3
MURF1 50 CANDVEQASNPLWQSRGSTTVSSGGRFRCPSCRHEVVLDRHGVYGLQRNL99
MURF2 50 CASDIFQASNPYLPTRGGTTVASGGRFRCPSCRHEVVLDRHGVYGLQRNL99
MURF3 47 CANDIFORANPYWTNRGGSVSMSGCRFRCPSCRHEVIMDRHGVYGLQRNL96
MURF1 100 L V E N I I D I Y K Q E S - S R P L H A K A E Q H L M C E E H E D E K I N I Y C L S C E V P T C S L 148

MURF2 100 L V E N I I D I Y K Q E S - T R P E - K K L D Q P M C E T H E E E R I N I Y C L N C E V P T C S L 146

MURF3 97 L V E N I I D I Y K Q E C S R P L Q - - K G S H P M C K E H E D E K I N I Y C L T C E V P T C S L 144
MURF1 149 CKVFGAHKDCEVAPL PTEYKROKSELSDGIAMLVAGNDRVQAVITQMEEV 198
MURF2 147 CKVPGAHKDCOVAPLTHVFQRQKSELSDGIAVLVAGNDRVQGVISOLEDT 196
MURF3 145 CKVFGAHQACEVAPLQSIFQGQKTELSNCISMLVAGNDRVQTIISQLEDS 194
MURF1 199 CQTIED NSRRQKQLLNORFETLCAVLEERKGELLQALAREQEEKLQRVRG 248
MURF2 197 CKTIEECCRKOKODLCEKFDHLYGILEERKTEMTQALTRTQEEKLEHVRT 246
MURF3 195 CRVTKENSHQVKEELSOKFDTLYAILDEKKSBLLORITQEGEKLGFIEA 244
MURF1 249 LIRQYGDELEGSSKLVESAIQSHEEPQHALYLOQAKELINEVGAMSKVEL 298
MURF2 247 LIBKYSDELENVSKLVESGIOFMDEPEMAVFLONAKTILQEIVEASKAFQ 296
MURF3 245 LILQYREQLEKSTKLVETAIOSLDEPGGATFLSSAKQLIKSIVEASKGCQ 294
MURF2 397 S S M P P A A D V L V T Q G E V V P I G S Q Q T T Q S E T S G P S A A E T A D P L F Y P S W Y K G Q 446
MURF1 356 E - - - - - - - L D V PE - - - - - - G S G - - - - - L H
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